## SEQUENCE LISTING

<110> Vreeland, Valerie Ng, Kwan L. The Regents of the University of California <120> Recombinant Vanadium Haloperoxidases and Their Uses <130> 023070 087100U5 <140> 09/151,189 <141> 1998-09-10 <160> 11 <170> PatentIn Ver. 2.0 <210> 1 <211> 2931 <212> DNA <213> Pucus distichus <220> <221> CD\$ <222> (228)..(2258) <223> vanadium bromoperoxidase <400> 1 cgcggacaag ccttggaaga gaggttgccc aattcaacag agcgaggccc gtgaaggtgt 60 ggaggacacg tgctacaagc tgatccacga gagcctcaac ttccctactg atacgggagt 120 ttgtactgcg ccgcgttgcc aaaaaccgca actttaaaca gcgctcgcga gcgccacatg 180 cttcccacgc atccacaaaa tcgacagtgg tatcgctgag cttgaat atg ctt tgc 236 Met Leu Cys cat gca gcg gac acg aca aga ggc tot cot atg cot gac acc gga gtg His Ala Ala Asp Thr Thr Arg Gly Ser Pro Met Pro Asp Thr Gly Val ctt cgg ttg ctc aca tca gag cag cgc gct aaa ggt tgg aga cgc cag Leu Arg Leu Leu Thr Ser Glu Gin Arg Ala Lys Gly Trp Arg Arg Gln tta gag ggg gag aaa tca cta ggt ttt cat cca agc gag acg cct tat 380 Leu Glu Gly Glu Lys Ser Leu Gly Phe His Pro Ser Glu Thr Pro Tyr 40 ate ang tac ttg gam ggc tet gag act tgg mag ang gtt amg ett eca 428 Ile Lys Tyr Leu Glu Gly Ser Glu Thr Trp Lys Lys Val Lys Leu Pro 55 acy gac ggc ata tog got too aag ato olg ggt aaa att atg goo agg 476 Thr Asp Gly Ile Ser Ala Ser Lys Ile Leu Gly Lys Ile Met Ala Arg gte ege ate get ace gee tig get gig gia etg gee gea eee igt tig Val Arg Ile Ala Thr Ala Leu Ala Val Leu Ala Ala Pro Cys Leu 90

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cac acc ggg gag gga aga cac ctc cag acc tgt aca aac tcc gac gat 620 His Thr Gly Glu Gly Arg His Leu Gln Thr Cys Thr Asn Ser Asp Asp 120 125 130
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tgc cas are act eac gga gaa act gat ttg gct acc atg ttc cac aag 764 Cys Cln Ile Thr Asn Gly Glu Thr Asp Leu Ala Thr Met Phe His Lys 165 170 175
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atc ctc gag gac tgc atc tta aac gga gat ttc agc att tgc gag gac 860 Ile Leu Glu Asp Cys Ile Leu Asn Gly Asp Phe Ser Ile Cys Glu Asp 200 205 210
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ttt gcc atc gac ata tcc ggt ccc gca ttc tcg gct acg aca ata ccc 956 Phe Ala Ile Asp Ile Ser Gly Pro Ala Phe Ser Ala Thr Thr Ile Pro 230 235 240
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gcc ggc ccc gaa gag tta gac gaa gag ctg cgt ttt acc cgt aac gcc - 138 Ala Gly Pro Glu Glu Leu Agg Dhe Ile Arg Asn Ala 375 - 380 - 385	88
cgc gac ctg gcc agg gtc tcc ttc gtg gac aat atc aac acc gaa gct 143 Arg Asp Leu Ala Arg Val Ser Phe Val Asp Asn Ile Asn Thr Glu Ala 390 395 400	6
tat ege ggg tet ett ate eta ett gag etg gga gee tte age agg eec 148 Tyr Arg Gly Ser Leu Ile Leu Leu Clu Leu Gly Ala Phe Ser Arg Pro 405 410 415	1
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acc tac ctt ctt cca caa gct atc caa gtg gga tcg cca acg cac cct 1820 Thr Tyr Leu Leu Pro Gln Ala IIe Cln Val Gly Ser Pro Thr His Pro 520 525 530	
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- Pro Cys Leu Ala Phe Asp Glu Val Thr Ala Ser Gly Val Phe Pro Glu 100 105 110
- Glu His Lys His Thr Gly Glu Gly Arg His Leu Gln Thr Cys Thr Asn 115 120 125
- Ser Asp Asp Ala Leu Asp Pro Thr Ala Pro Asn Arg Arg Asp Asn Val
- Ala Phe Ala Ser Arg Arg Asp Ala Ala Arg Arg Glu Arg Asp Gly Thr 145 150 155
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- Cys Glu Asp Val Pro Ala Gly Asp Pro Ala Gly Arg Leu Val Asn Pro 210 215 220
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